5 (Amended). An oligonucleotide characterized in that the gene sequence of a spacer region between a gene coding 16S rRNA and a gene coding 23S rRNA of *Pectinatus frisingensis* has at least one of the following sequence group or the corresponding complementary sequence:

5'-CCATCCTCTTGAAAATCTC-3' ① (SEQ ID NO:5)

5'-TCTCRTCTCACAAGTTTGGC-3' ② (SEQ ID NO:6).

6 (Amended). An oligonucleotide characterized in that the gene sequence of a spacer region between a gene coding 16S rRNA and a gene coding 23S rRNA of *Pectinatus cerevisiiphilus* has at least one of the following sequence group or the corresponding complementary sequence:

5'-CACTCTTACAAGTATCTAC-3' ③ (SEQ ID NO:7)

5'-CCACAATATTTCCGACCAGC-3' ④ (SEQ ID NO:8)

5'-AGTCTTCTCTACTGCCATGC-3' ⑤ (SEQ ID NO:9).

11 (Amended). A method as claimed in claim 9, wherein the nucleotide sequence coding the 16S rRNA gene of *Pectinatus frisingensis* has the following sequence:

5'-CGTATCCAGAGATGGATATT-3' ⑥ (SEQ ID NO:10)

12 (Amended). A method as claimed in claim 10, wherein the nucleotide sequence coding the 16S rRNA gene of *Pectinatus cerevisiiphilus* has the following sequence:

5'-CGTATGCAGAGATGCATATT-3' ⑦ (SEQ ID NO:11)